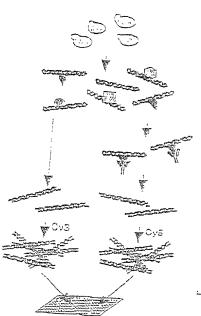
Title: Genome-Wide Location and Function...

Inventors: John Wyrick, et al.



Crosslink protein to DNA In vivo with formaldenyae

Breek open calls and snear DNA

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Reverse-crossinks, blum DNA and ligate to unidirectional linkers

LIVI-FOR

Hypridize to erray

FIGURE 1

Title: Genome-Wide Location and Function...
Inventors: John Wyrick, et al.

Triplicate experiments	Scanning	Single-array error model	Weighted average
ChiP-1	Scan-:	- Data-1	
CnIF-2	Scan-2	- Date-2	Final data
ChiP-S	Scan-3	- Data-8	

FIGURE 2

General Transcription Factors SWI/SNF Complex SAGA

ORC complex DNA Polymerase

SRB/Medialor Complex

RNA Polymerase II

Interests:

Title: Genome-Wide Location and Function...

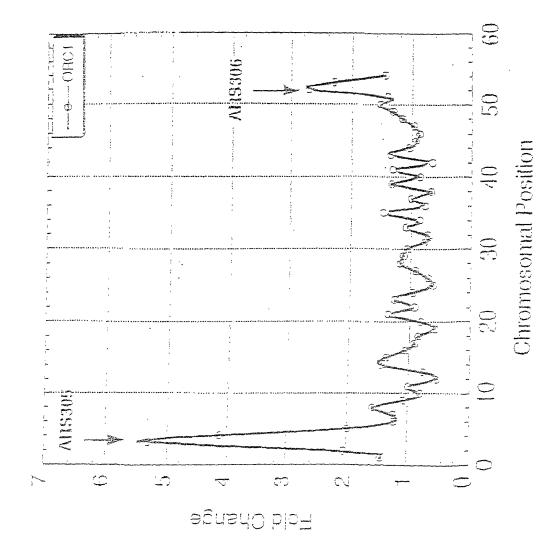


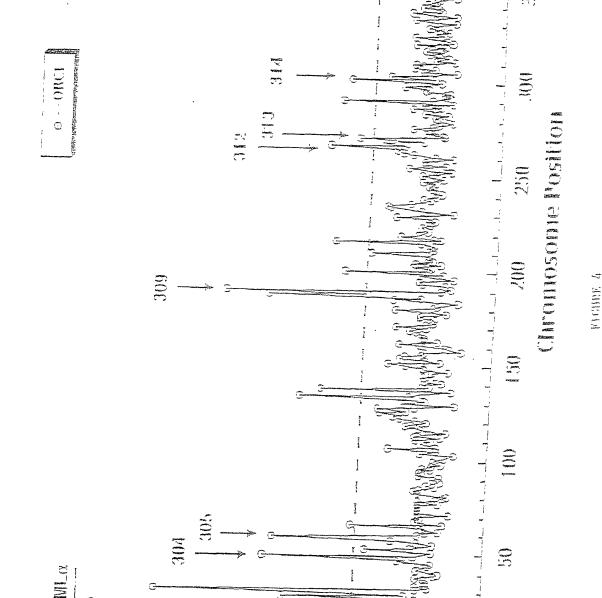
FIGURE 3

elomeric repeats

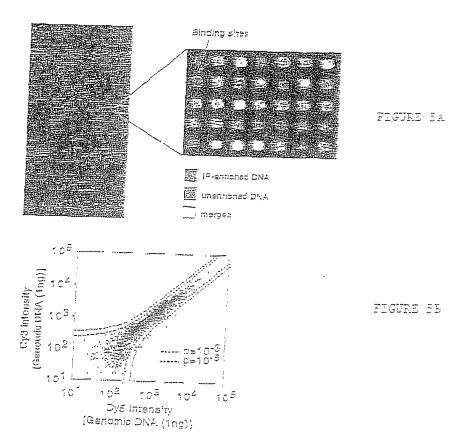
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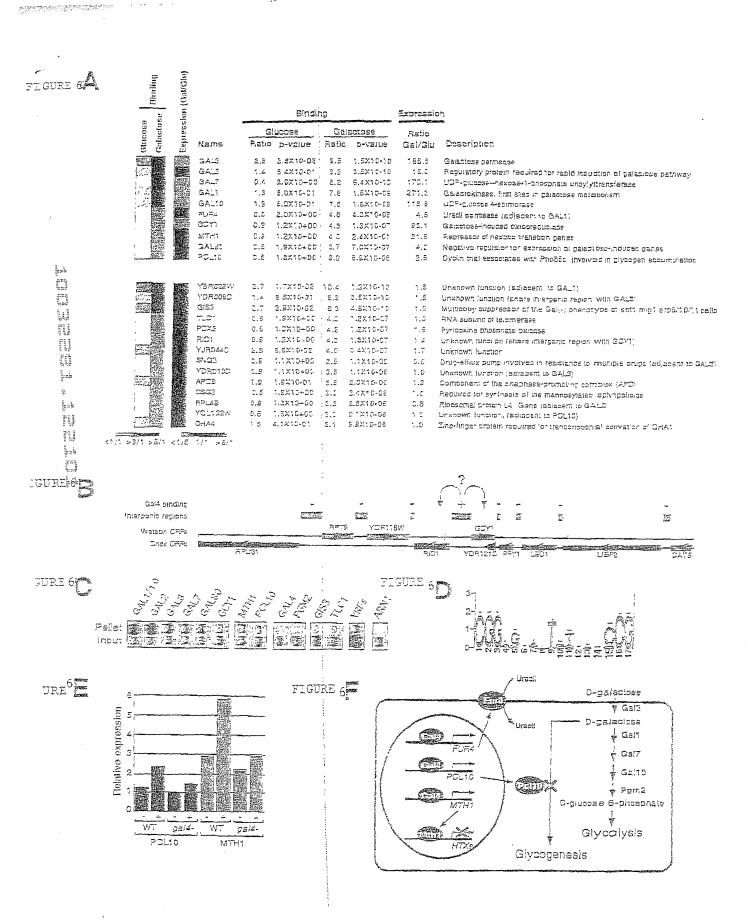
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Title: Genome-Wide Location and Function...



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Docket No.: 0399.1212-005
Title: Genome-Wide Location and Function...

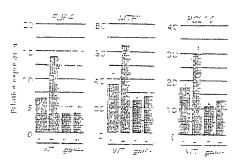


FIGURE 6G

Title: Genome-Wide Location and Function...

٠	FIGUR	E 7					
	Sinding	Expression	Name	Binding			
	2	4 1 t		before		after	
	belore alter	25.55 25.55 25.55 25.55 25.55 25.55 25.55 25.55 25.55 25.55 25.55 25.55 25.55 25.55 25.55 25.55 25.55 25.55 25		rat,o p-valu	e rati	o p-valu	e Description
			PRM:	2.7 1.95-04	6.2		
			ERG24 PO_2	2,7 1,8E-04	6,2		
			F132	8.6 4.2E-CS 2.6 2.2E-04	5,8 2,6		
			STE12	4 1 1.4E-08	4.4	3,15-04	
· .			FUS1	5.0 2.8E-07	4	5.52-04	
h=h			CDC48	2.0 1.55-04	4.5	1,55-00	hadusted for cell division and nometypic memorana tuaion
7.17 2.27			Y0R0270	2.4 5.8E-04	3.2	1,5E-03	
April 1980			FUSS	3.5 3.8E-C5	€,4		
1:1			PEP; MET2	3.2 3.9 <b>5</b> .03 2.1 4.85.00	€,∠ Ω.&	2.0E-03 3 2E-03	
क्षित्रेष्ट सम्बद्ध			YOR1285	2.1 4.8E-00 0.7 1.0E-00	5 2	1.0E-04	
Harry Brand	洒鬱		UTHI	1.7 2 8E-02	5.2	1.05-04	
			AFR'	1.0 8,68-0:	٤.٠	1.0E-04	
I >5/1 B			3102	1 4 1.1E-01	€.0	: EE-04	
jek 📗			YNR046W	1.9 7.6E+03	4.2	1.4E-04	
2			MSS11 YOR3430	1.0 E.1E-01	۵,9	1.5E-04	
4/1			TURISHEL THS:	" 1 7.5E-0" 0 9 1.0E-00	5.2 4,1	4,18404 4,25-04	
g=4 <del>4+</del> /) g			SCHS	1.3 4.1 <b>E</b> -01	4.2	4.55-04	
			KAR5	1,2 3,7E-01	3.8	4.7E-04	Memorana protein racultas ter nemetypio hubigar fusion
A line lines			YER046W-#		3.8	5.1E-04	Protein of unknown junction
3/1			YPL1920	0.9 1.0E-00	3.5	5.5E+04	Process of unknown function
			SPC25 YERD197	1.9 2.05-01 1.3 2.05-07	47	6 6E-04 8,8E-04	Protein of the spingle pole body  Moderately Similaric mammalish neutral sphingon velimesso
A A S 2/			Y!L053C	1.0 8.7E-01	4.† 3 E	7.15.04	Protein of unknown function
2/1 🕅			PGIM*	2,0 0,1≅-00	3,8	7.25-04	Phosphogiupomulase
			YILDS70	1.1 5.5E-D1	3.4	E.2E-04	Ризтел ст инклема тупског
1.			Y"L1980	1.8 7. <b>35.</b> 52	úΞ	1.25-03	
<1:1 -		50170	AGA1	1.4 1.1E-01	3 1	1,95-03	
< i . 1 =			YOL1550 VT:1	1.0 7.3E-21	3,0	215.03	Similar to 3, carevisiae giyoch 1,4-aip ha-giyoosidaae Synabiobravin (v-SNARE, homolog
			DIKT	0.5 1 0E-00 0.8 1,02+00	3 C 3.D		tunothed to setudie solustrou sug Kalinodemi.
			FARS	1 3 2.25-01	3.0		involved in call cycle arrest for mating
			PHCS	1.0 8.3E-0:	2.5	2.75.03	CDK inhibitor for Pho80p-Pho85p com piex
			YNAD54C	1.5 7 4E-02	2.7	4.8E-03	Similar to Rhocopacier capsulatus both C protein
			HYM1 FIG1	1.85-01	2.7	4,95-00	Protein with similarity to Aspergillus nudulans nym4
, met	·ene	200	(12)	1.2 8.8E-0°	2.9	5 DE-03	Protein required for efficient mating
>10/1 影響的		•					
5/1 🗓	स्थानका क		DM IC		, ,		The second of th
			BMH2 YGR1880	2,1 3.2E-03 1,7 2.4E-03	4.5 4.5		Homolog to 14-3-3 protein; sojacent to Ty element YDROdelleS Cell wall protein; sdjacent to Ty element YGRWoelleS*
			YNADESC	1.9 5.7E-C3	4.3		Unknown function; adjacent to Ty element YNRWeigme4
2/7 麗			MSC1	1.9 1,82-02	4.3	2.9E-04	Municopy suppressor of sect; solatent to Ty element YNPWs:gma4
			TYET	1.0 1.0E-00	3.3	4.62-94	Basio helix-loop-nailx transcription factor
			YN9051C	1 B 1.4E-02	3.8	4.7E-04	Similar to onicken nucleolin; adjacent to Ty element YNEW/sigma-
1:1 🖁			YNR048W	1.9 9.25-03	3.8		Protein of unknown junction; adjacent to Ty element YNRWsigma4
	Training B		YGL050W YIL035W	1.7 2.0E+02 7.1 6.5E-01	3.8 3.4		Protein of unknown function; adjecent to Ty element YGLOsigme? Similar to Mel4e and to CRES proteins; a djacent to YILOS7O
1/0 配			YNE0570	2.0 1.3E-02	3.4		Smilla. To Mere and to Greek proteins; a dijadent to 11605/0 Protein of unknown tunction
7/4 間	7700 <b>8</b>		SDL1	1.6 7.25.CZ	22		Senne penydratase; adjacent to YIL189C
	: E		YDR2810	1.2 4.2E-01	3,1		Protein of unknown function; adjacent to Ty element YDAWsigmas
			YDA100W	1.1 3.8≅.01	5,5		Protein of unknown function; adjacent to Ty siement YDRCdeltaß
1/5			YNL0430	1.1 5.7E-01	3.D 3.D		Protein of unknown function; adjacent to Ty element YNLWsigma2
			YDL154W YHB1	1.0 7.9E-01 7.0 8.3E-01	2.3		Protein with similarity to Zinc metalloprot sinesest adjacent to YOL155 Flevohemoglobin of unknown function; adjacent to PHOS1
1:1 1/2 1/5 <1/10			PANS	1.8 B.SE-02	4.5		Companent of Pab1p-atimulated poly(A)   Fibonuclasse
-1110 (20)			CWP2	1.8 1,5E-02	2.5	3.8E-03	major constituent of the call wall containing GPI-anchor
	MARKET E		YKL097C	1.8 1.8%-02	2.5	3.8E-03	Protein of unknown function; adjacent to CWP2

Title: Genome-Wide Location and Function...

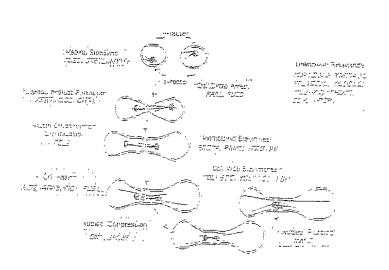
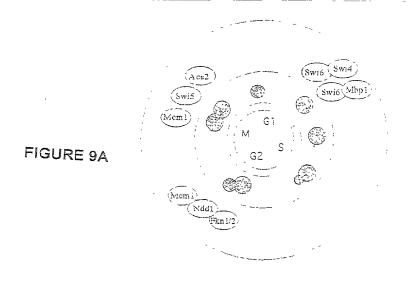
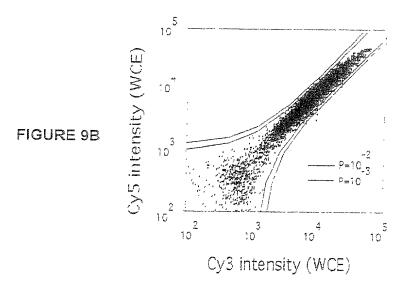
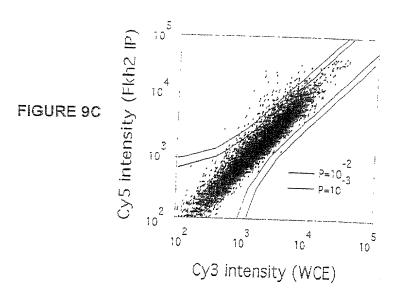


FIGURE 8

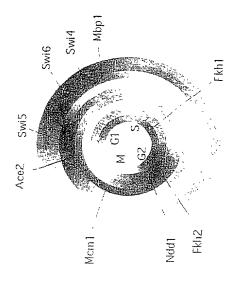
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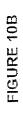


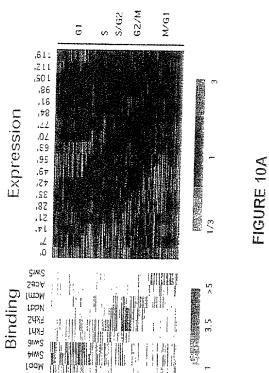




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Inventors: John Wyrick, et al.



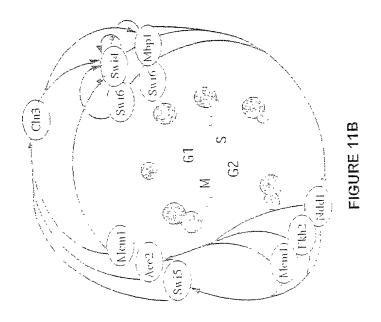


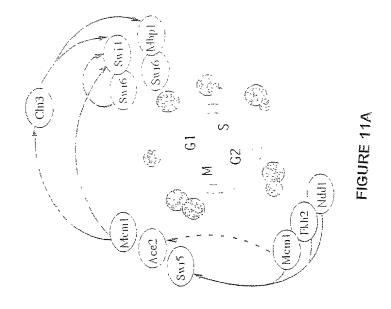


Docket No.: 0399.1212-005

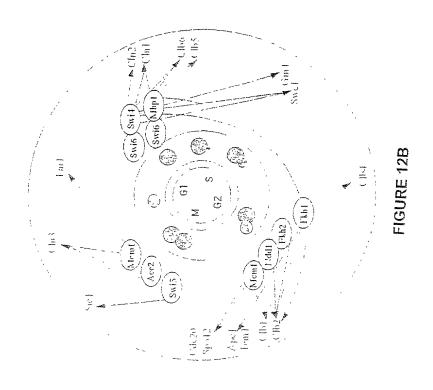
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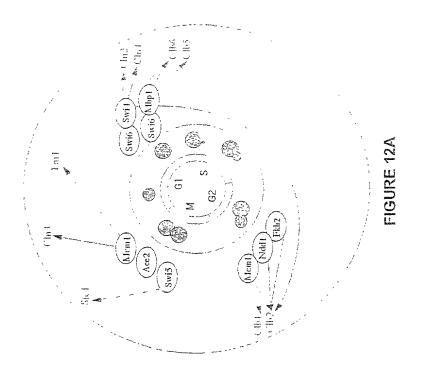
Inventors: John Wyrick, et al.





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Title: Genome-Wide Location and Function...
Inventors: John Wyrick, et al.





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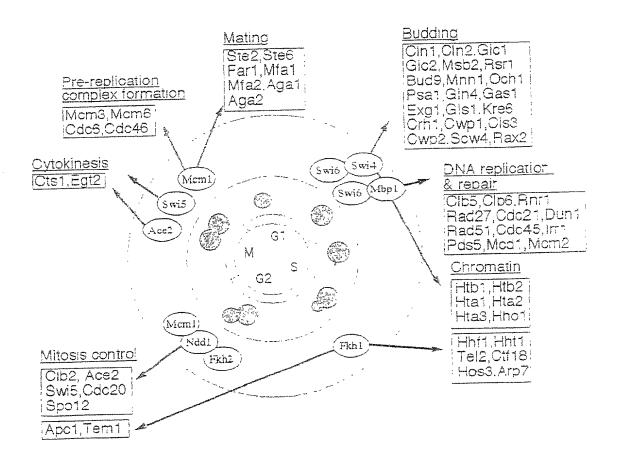


FIGURE 13

Title: Genome-Wide Location and Function...

